

FIG.1

(2 of 90)

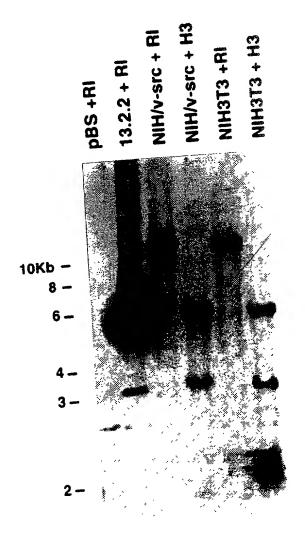


FIG.2A

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#### 322 cDNA (5.4Kb)

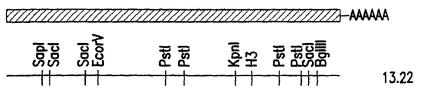


FIG.2B

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ç	ggaa gtte ggca	gtca	agca	gac	ctac	gaç	gaaç	gto	gga	gct	gcc:	ttt	gga	aga	cca	ggt:	tgg	tga	cct	gga	60 120 180 2
.81 a	agce	cca	ccaa	agaa	agti	tgt1	tgca	aga	ggt	cca	cgt	gag	cac	cgt	gga	gaa	gac	aga	gga	gga	240
	A	H	Q	E	V	V	A	E	V	E	V	S	T	V	E	K	T	E	E	E	22
241 (	gca	ggg:	agga	agga	agga	agaç	ggc1	tga:	agg	ggg	cgt	ggt	ggt	aga	agg	aac	agg	aga	atc	ctt	300
23	Q	<u>G</u>	G	G	G	E	A	E	G	<u>G</u>	V	V	V	E	G	T	G	E	S	L	42
301 ( 43	gcc P	CCC	tgas E	gaaa K	act:	ggc† A	tgaq E	gcc P	cca Q	gga E	ggt V	ccc P	cca Q	gga E	agc A	tga E	gcc P	tgc A	tga E	gga E	360 62
361 g	gct:	gat	gaa!	gagı	caga	aga:	gate	gtg	tgt	ctc	tgg	agg	aga	cca	cac	tca	act	gac	aga	cct	420
63		M	K	S	R	E	M	C	V	E	G	G	D	H	T	Q	L	T	D	L	82
121	aag	tcc		aga	gaa:	gacı	gct:	gcc	caa	aca	ccc	aga	agg	cat	tgt	cag	tga	ggt	gga	gat	480
83	S	P		E	K	T	L	P	K	H	P	E	G	I	V	S	E	V	E	M	102
181	gct	gtc	ctc	tca	gga	aag	aat	caa	ggt	aca	ggg	aag	tcc	ctt	gaa	gaa	act	ctt	cag	tag	540
103	L	S	S	Q	E	R	I	K	V	Q	G	S	P	L	K	K	L	F	S	S	122
541	ctc	agg	ctt	aaa;	gaa	gct	gtc	tgg	gaa	gaa	gca	gaa	ggg	ıgaa	iacg	agg	agg	tgg	999	aga	600
123	S	G	L	K	K	L	S	G	K	K	Q	K	G	K	R	<u>G</u>	G	G	<u>G</u>	D	142
501	cga	aga	-	tgg	aga	ata	cca	aca	cat	tca	cac	cga	atc	ccc	aga	gag	tgc	tga	tga	gca	660
143	E	E		G	E	Y	Q	H	I	H	T	E	S	P	E	S	A	D	E	Q	162

FIG.3A

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661	gaagggagagagetetgegtegteeceegaggageetgaggagaeeaegtgtetggagaa	720
163	KGESSASSPEEPEETTCLEK	182
721 183	agggccgctggaagcacccaggatggggaagctgaggaagga	780 202
781	gaagaggaaggatcactccctgggcatccttcaaaaagatggtgacacccaagaaacggt	840
203	KRKDHSLGILQKDGDTQETV	222
841	ccgaagaccttctgagagtgacaaggaggaagagctggagaaggtcaagagcgccacctt	900
223	R R P S E S D K E E E L E K V K S A T L	242
901	gtcctccactgatagcacagtgtcagaaatgcaagatgaagtcaaaactgttggtgagga	960
243	SSTDSTVSEMQDEVKTVGEE	262
961	acaaaagccagaggaaccaaagcgtagggtggatacttcagtgtcttgggaagcactgat	1020
263	Q K P E E P K R R V D T S V S W E A L I	282
1021 283	ttgtgtcggatcatccaagaagaggcaaggaaggcatcctcttcagatataagagggccCVGSSK <u>KRAR</u> KASSSDIRGP	1080 302
1081	aaggacactgggaggggacagtcacagagcagagggggccagcaaagacaaagaagccg	1140
303	RTL <u>GGGQSOSRGG</u> QQRQRSR	322
1141	aacagacgctgttcctgccagcacccaggagcaggaccaagcgcaaggaagttcctcacc	1200
323	T D A V P A S T Q E Q D Q A Q G S S S P	342
.201	cgagccagcgggaagcccttccgaaggggaaggtgtctccacttgggagtcatttaaaag	1260
343	EPAGSPSEGEGVSTWESFKR	362

FIG.3B

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1261 363	attagtcactccaagaaaaaaatccaagtcaaaactggaagaaaga	1320 382
1321	tctagttgtaggagcaggttgtccactgagatcgaaccgtgtagagaagaatcttgggtt	1380
383	L V V G A G C P L R S N R V E K N L G F	402
1381	tccattaagaaattcatccccggacggcggaagaaaagggcagatgggaaggcaagaaca	1440
403	PLRNSSPDGGRKGQMGRQEQ	422
1441	agccactgtggaagactcagggccagtggagataaatgaggacgagcctgatgtcccagc	1500
423	A T V E D S G P V E I N E D E P D V P A	442
1501	agtcgtgcctctgtctgagtatgatgcagtggagagggagaagatggaagcccaggggaa	1560
443	V V P L S E Y D A V E R E K M E A Q G N	462
1561	tgcggagctgcccagctgctggggctgtgtagtgtccgaggagctcagtaagactctggt	1620
463	A E L P S C W G C V V S E E L S K T L V	482
1621	ccacactgtgagtgtcgcagtcattgatgggaccagggcagtcaccagtgtcgaagagcg	1680
483	H T V S V A V I D G T R A K T S K E E R	502
1681	gteteettegtggatateegetteegtaacagaacetettgaacacacagegggagaage	1740
503	SPSWISASVTEPLEHTAGEA	522
1741	catgccacctgttgaagaggtcactgaaaaagacatcattgcagaagaaactcctgtgct	1800
523	M P P V E E V T E K D I I A E E T P V L	542
1801	cacccagacgttaccagagggtaaagatgcccatgacgacatggtcaccagtgaagtgga	1860
543	TQTLPEGKDAHDDMVTSEVD	562

FIG.3C

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1861	tttcacetcagaagetgtgacagecacagaggacetcagaggeteteegtaetgaagaagt	1920
563	FTSEAVTATETSEALRTEEV	582
1921	taccgaagcatcgggggccgaagagaccacagacatggtgtccgcagtttcccagctgac	1980
583	T E A S G A E E T T D M V S A V S Q L T	602
1981 603	tgactccccagacaccacagaggaagccaccccagttcaggaggtagagggtggtgtgct D S P D T T E E A T P V Q E V E G G V L	2040 622
2041	agatacagaagaagaggagcgcagacgcaggccatcctccaagccgttgcagacaaggt	2100
623	DTEEEERQTQAILQAVADKV	642
2101	gaaagaggagtcccaggtgcctgcaacccagactgtgcagagaacggggtcaaaagcact	2160
643	KEESQVPATQTVQRTGSKAL	662
2161	ggagaaggttgaggaggtagaggagtcccgaagtgctggcttcggagaaagagaagga	2220
663	EKVEEVEDSEVLASEKED	682
2221	cgttatgccgaaaggacccgtgcaggaagctggagctgagcatcttgcacagggctctga	2280
683	V M P K G P V Q E A G A E H L A Q G S E	702
2281	gactggacaggctactccagagagccttgaagttcctgaagtcacagcagatgtagacca	2340
703	T G Q A T P E S L E V P E V T A D V D H	722
2341	tgtcgccacgtgccaggttatcaagctccagcagctgatggaacaggccgtggcccctga	2400
723	V A T C Q V I K L Q Q L M E Q A V A P E	742
2401	gtcatccgaaaccttgacagacagtgagacaaatggaagcactcccttagcagattcaga	2460
743	SSETLTDSETNGSTRLADSD	762

FIG.3D

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2461	cactgcagatgggacacagcaagatgaaaccattgacagccaggacagtaaagccactgc	2520
763	T A D G T Q Q D E T I D S Q D S K A T A	782
2521	agctgtcaggcagtcacaggtcacagaagaagaggcggctactgctcagaaagaggagcc	2580
783	AVEQSQVTEEEAATAQKEEP	802
2581	ttcgacactacctaataatgttccagcccaggaagaacatggggaagaaccaggaagaga	2640
803	S T L P N N V P A Q E E H G E E P G R D	822
2641	tgttcttgaacctacacagcaagagcttgctgctgcagccgtgcccgtctggcaaaagac	2700
823	V L E P T Q Q E L A A A V P V W Q K T	842
2701 843	tgaggtgggtcaagagggtgaggttgactggttggatgga	2760 862
2761	ggaggtgtttgtacactctggacccaacagtcaaaaggctgctgatgtgacatatgacag	2820
863	EVFVHSGPNSQKAADVTYDS	882
2821	tgaagtgatgggagtggccgggtgtcaggaaaaggagagtactgaagtgcagagtcttag	2880
883	EVMGVAGCQEKESTEVQSLS	902
2882	cctggaggagggagagatggaaactgacgttgaaaaggagaaaagggagacaaagccaga	2940
903	L E E G E M E T D V E K E K R E T K P E	922
2941	gcaagtgagtgaagaaggtgagcaggaaacagccgctcctgagcatgaaaggaactacgg	3000
923	Q V S E E G E Q E T A A P E H E R N Y G	942
3001	gaagccagtcctgacacttgacatgcccagctcagagagggggaaggcactgggaagcct	3060
943	K P V L T L D M P S S E R G K A L G S L	962

FIG.3E

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3061	tggaggaagcccttctctcccagaccaagacaagcaggttgcatagaggttcaagttca	3120
963	GGSPSLPDQDKAGCIEVQVQ	982
3121	aagcctggacacaacagtcactcaaacagcagaagctgtggaaaaggtcatagaaacggt	3180
983	S L D T T V T Q T A E A V E K V I E T V	1002
3181 1003	tgtgatttcagagacaggtgaaagtccagagtgtgtaggtgcacacttattaccagctga V I S E T G E S P E <u>C</u> V G A <u>H</u> L L P A E —Zn-finger—	3240 1002
3241	gaagtcctctgcaacgggtggccactggactcttcagcatgcagaggacacggtacccct	3300
1023	K S S A T G G H W T L Q H A E D T V P L	1042
3301	ggggcctgagtctcaggcagaatccatcccaatcatagtaactcctgctcctgaaagcac	3360
1043	G P E S Q A E S I P I I V T P A P E S T	1062
3361 1063	cctacatcctgacctacaaggagaaataagcgcatcccagagagag	3420 1082
3421	ggacaagccagatgctggtcctgatgctgacggcaaggaggagtacagcaatcgacaaagt	3480
1083	D K P D A G P D A D G K E S T A I D K V	1102
3481	cctcaaggctgaacctgagatcctggaacttgagagtaagagcaacaagattgtgctgaa	3540
1103	L K A E P E I L E L E S K S N K I V L N	1122
3541	cgtcattcagacagccgttgaccagttcgcacgtacagaaacagcccccgaaactcatgc	3600
1123	V I Q T A V D Q F A R T E T A P E T H A	1142
3601	ttatgattcacagacccaggttcctgcaatgcgcttggacagcagggagcccaacagatg	3660
1143	Y D S Q T Q V P A M R L D S R E P N R C	1162

FIG.3F

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3661 1163	ctgga W T																			3720 1182
3721 1183	gcaag Q V	tcct L	-															tgc A		3780 1202
3781 1203	tgaaa E S	igcgc A																	aaa K	3840 1222
3841 1223	ggagd E H	atgo ł A																		3900 1242
3901 1243	aaacc N L																	atg C		3960 1262
3961 1263	ccaaa Q K	iagtt ( L	gag R	gtc S	cagg R	gaag K	gaaq K	gaaa K	atgi C	tcta L	acca P	aagi S	tca Q	gtc S	aaa K	gag. R	aac T	aag R	gcc P	4020 1282
4021 1283	caggo R Q																			4080 1302
4081 1303	tgctc A H					aaga R												gct: L		4140 1322
4141 1323	gttgg L G																	tcca P		4200 1342
4201 1343	gattt I S	ccac			gago			gaca									aga	gcca	agc	4260 1346

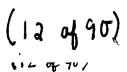
FIG.3G

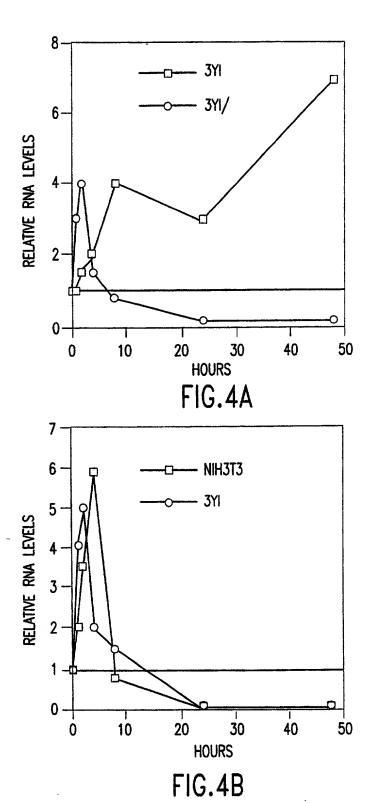
# (11 of 90)

4261 4321 4381	taacatttcctcgtttcaagactgcctttgatttgccccttgatgccgtccgt	4320 4380 4440
4441	${\tt acctattgtatatttttttctaacgtttaagcacatgctttttgtattatgcaatatata}$	4500
4501	acgggtgtgcagccatagcgacgctttgaaaagctccaagcctcaactgtaacctgcagc	4560
4561	aaacagataacattcctggcaagaagagacaagtctttttaaaagtttactgatgcttag	4620
4621	atctgtgggcttctagtcctctgaaagtggttgttttcctatgcacagcgagctcag <u>aaa</u>	4680
4681	<u>taaa</u> aaccccattttgaaacatccaggatgtcccaatattaccatgattttttcccccct	4720
4721	ttttgctaatccagtccaggttggaaagaagtctcctctgtgtcagattaagccctgtct	4780
4781	cttaatgatatggacaaatgagtgtgcctaaggccatgagatgtttcctaatgcagaagg	4840
4741	aatctgttgtacgtttttttgattgtactcttctatgctggaccgaattcatatgcagat	4900
4901	cgaagtgagtcctgttctttacagatggtattttgatagata	4960
4961	atatctgtgccccttctttaagaacaatgttgcattatgttcctttggataaattgtgat	5020
5021	ttgacaactgattta <u>aataaa</u> catatttgactac(A).	

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FIG.3H





rat-6/mos rat-6/src rat-6/myc rat-6/ras rat-6/raf-1 rat-6

FIG.5

pBluescript KS II + T7/luciferase + 13.2.2 c.c. +endo F + 13.2.2 c.c. + 13.2.2 + Smal + T3/luciferase + 13.2.2 c.c. +

T7 RNA Pol T3 RNA Pol

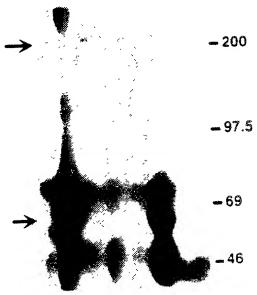


FIG.6

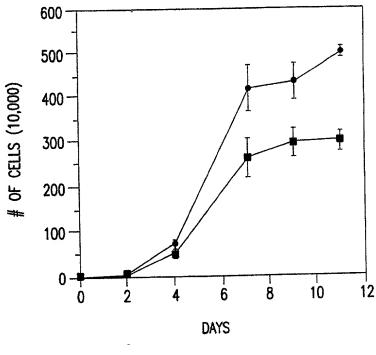


FIG.7A

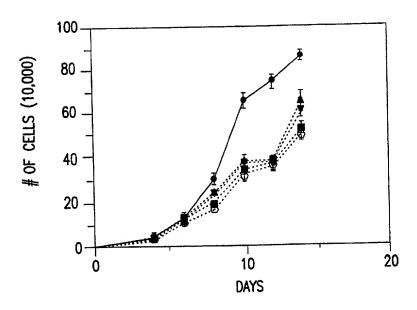


FIG.7B

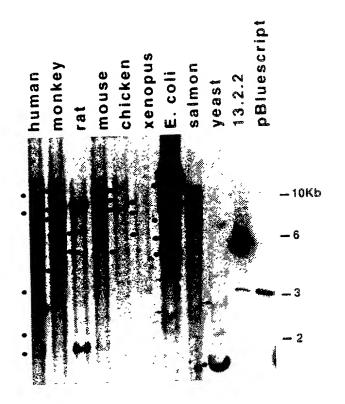
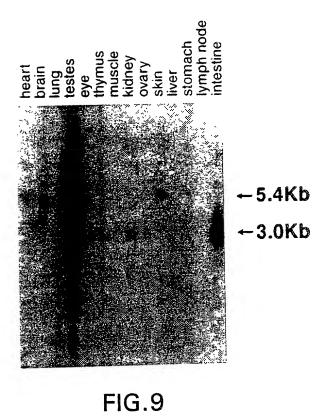


FIG.8



(19 of 90) M G A G S S T E Q R S P E Q P A G GAC ACG CCG AGC GAG CTG GTG CTC AGT GGC CAT GGG CCC GCA GCT GAA GCC TCG D T P S E L V L S G H G P A A E A S G A A G D P A D A D P A T K L P O K AAT GGC CAG CTG TCT TCT GTC AAC GGC GTA GCT GAA CAA GGA GAT GTC CAT GTC N G Q L S S V N G V A E Q G D V H V Q E E N Q E G Q E E E V V D E D V G CAG CGA GAG TCA GAA GAT GTG AGA GAA AAA GAC CGA GTT GAA GAA ATG GCG GCC O R E S E D V R E K D R V E E M A A AAC TCC ACA GCT GTT GAA GAT ATC ACA AAG GAT GGG CAG GAG GAG ACA TCA GAA N S T A V E D I T K D G Q E E T S E ATA ATT GAA CAG ATC CCT GCT TCA GAA AAC AAT GTG GAA GAA ATG GTA CAG CCT I I E Q I P A S E N N V E E M V Q P

FIG.11A

(20 of 90) 450 468 459 GCT GAG TCC CAG GCT AAT GAT GTT GGC TTC AAG AAA GTA TTT AAA TTT GTT GGT A E S Q A N D V G F K K V F K F V G 504 513 522 495 TTT AAA TTC ACG GTG AAG AAG GAT AAA AAT GAA AAG TCA GAT ACT GTC CAA CTA F K F T V K K D K N E K S D T V Q L 576 585 567 558 CTC ACT GTC AAG AAG GAT GAA GGC GAA GGG GCA GAA GCC TCT GTC GGA GCT GGA L T V K K D E G E G A E A S V G A G 621 630 639 612 GAC CAC CAG GAG CCC AGT GTG GAG ACT GCC GTC GGA GAG TCA GCA TCC AAA GAA D H Q E P S V E T A V G E S A S K E 693 675 684 666 AGT GAG CTG AAG CAA TCC ACA GAG AAG CAA GAA GGC ACC CTG AAG CAA GAA CAG S E L K Q S T E K Q E G T L K Q E Q 729 738 720 711 AGC AGC ACA GAA ATC CCC CTT CAA GCC GAA TCT GAT CAA GCG GCT GAG GAA GAA S S T E I P L Q A E S D Q A A E E E 774 783 792 801 A K D E G E E K Q E K E P T K S P E 837 846 855 828 TCC CCG AGC AGC CCA GTC AAC AGT GAG ACA ACA TCT TCC TTC AAG AAG TTC TTC P S S P V N S E T T S S F K K F F

**FIG.11B** 

( 21 of 90) ACT CAC GGT TGG GCC GGC TGG CGC AAG AAG ACC AGC TTC AAG AAA TCA AAA GAG THGWAGWRKKTSFKKSKE GAT GAT CTG GAA ACT GCC GAG AAG AGA AAG GAG CAA GAG GCA GAA AAA GTA GAC D D L E T A E K R K E Q E A E K V D GAG GAA GAA AAG GAA AAG ACA GAG CCA GCC TCG GAG GAG CAG GAG CCG GCA GAA EEEKEKTEPASEEQEPAE GAC ACA GAC CAG GCC AGG TTG TCA GCA GAC TAC GAG AAG GTG GAG CTG CCT TTG D T D Q A R L S A D Y E K V E L P L GAA GAC CAG GTT GGT GAC CTG GAG GCA TCG TCA GAG GAG AAG TGT GCT CCT TTG E D Q V G D L E A S S E E K C A P L GCA ACG GAA GTG TTT GAT GAG AAG ATG GAA GCC CAC CAA GAA GTT GTT GCA GAG ATEVFDEKMEAHQEVVAE V H V S T V E K T E E E Q G G G E GCT GAA GGG GGC GTG GTA GAA GGA ACA GGA GAA TCC TTG CCC CCT GAG AAA A E G G V V V E G T G E S L P P E K

**FIG.11C** 

( 22 of 90) 1314 1332 1341 CTG GCT GAG CCC CAG GAG GTC CCC CAG GAA GCT GAG CCT GCT GAG GAG CTG ATG LAEPQEVPQEAEPAEELM 1359 1377 1386 1368 1395 AAG AGC AGA GAG ATG TGT GTC TCT GGA GGA GAC CAC ACT CAA CTG ACA GAC CTA K S R E M C V S G G D H T Q L T D L 1413 1422 1431 1440 1449 AGT CCT GAA GAG AAG ACG CTG CCC AAA CAC CCA GAA GGC ATT GTC AGT GAG GTG SPEEKTLPKHPEGIVSEV 1467 1476 1485 1494 GAG ATG CTG TCC TCT CAG GAA AGA ATC AAG GTA CAG GGA AGT CCC TTG AAG AAA E M L S S Q E R I K V Q G S P L K K 1530 1539 1548 1557 1566 CTC TTC AGT AGC TCA GGC TTA AAG AAG CTG TCT GGG AAG AAG CAG AAG GGG AAA L F S S G <sup>-</sup> L K K L S G K K Q K G K 1575 1584 1593 1602 1611 CGA GGA GGT GGG GGA GAC GAA GAG CCT GGA GAA TAC CAA CAC ATT CAC ACC GAA RGGGGDEEPGEYQHIHTE 1629 1638 1647 1656 1665 TCC CCA GAG AGT GCT GAT GAG CAG AAG GGA GAG AGC TCT GCG TCC CCC GAG SPESADEQKGESSASSPE 1683 1692 1701 1710 1719 GAG CCT GAG GAG ACC ACG TGT CTG GAG AAA GGG CCG CTG GAA GCA CCC CAG GAT E P E E T T C L E K G P L E A P Q D

FIG.11D

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GGG		1737 GCT						1755 TCC						1773 AGA			1782 ATC
G	E	A	 Е	E	G	T	T	 S	D	G	E	 К	K	R	<b>Е</b>	G	Ī
ACT		1791 TGG	GCA		1800 TTC			1809 ATG			1818 CCC			1827 CGG			1836 AGA
T	Р	W	Α	S	F	K	K	M	٧	T	Р	K	K	R	٧	R	R
CCT		1845 GAG	AGT		1854 AAG			1863 GAG				GTC		1881 AGC	GCC		1890 TTG
Р	S	F	S	D	K	E	E	E	L	E	K	٧	K	S	Α	Ţ	L
TCC		1899 ACT			L908 ACA							GAA			ACT		1944 GGT
S	S	Τ	D	S	T	٧	S	E	М	Q	D	E	٧	K	Т	٧	G
GAG		1953 CAA	AAG		1962 GAG		CCA			AGG		GAT		1589 TCA	GTG		1998 TGG
	GAA			CCA	GAG	GAA	CCA	AAG	CGT	AGG	GTG	GAT	ACT	TCA		TCT	
E	GAA E	CAA	K	CCA P	GAG E E 2016	GAA  E	CCA P	AAG  K 2025	CGT  R	AGG  R	GTG V 2034	GAT D	ACT T	TCA S 2043	V	TCT S	TGG W
E GAA	GAA E	CAA Q 2007 CTG	K ATT	CCA P	GAG E E 2016	GAA E GGA	P Z	AAG  K 2025	CGT R R	AGG R AAG	V 2034 AGA	GAT D GCA	ACT T	TCA S S 2043 AAG	V	TCT S	TGG W
E GAA  E	GAA E GCA A	CAA Q 2007 CTG	K ATT I	CCA P TGT C	GAG E 2016 GTC V 2070	GAA E GGA  G	CCA P  TCA S	AAG  K 2025 TCC  S	CGT R R AAG  K	AGG R R AAG K	GTG  V 2034 AGA  R	GAT D GCA A	ACT T T AGG  R	TCA S 2043 AAG K	V GCA	TCT S TCC S	TGG W 2052 TCT S
E GAA  E	GAA E GCA A	CAA Q 2007 CTG L	ATT I GAA	CCA P TGT C GGA	GAG E 2016 GTC V 2070 GGG	GAA E GGA  G	CCA P TCA S AGG	AAG  K 2025 TCC  S	CGT R AAG K CTG	AGG R AAG K K GGA	GTG V 2034 AGA R R 2088 GGG	GAT D GCA A GAC	ACT T AGG R	TCA S 2043 AAG K 2097 CAC	V GCA	TCT S TCC S	TGG W 2052 TCT S 2106 GAG
GAA E TCA S	GAA E GCA A GAT D	CAA Q 2007 CTG  L 2061 GAT	ATT I GAA E	CCA P TGT C GGA G	GAG  E 2016 GTC  V 2070 GGG  G	GAA E GGA G CCA P	P Z Z AGG R	AAG  K 2025 TCC  S 2079 ACA  T	CGT R AAG  K CTG	AGG R AAG K K GGA G	GTG V 2034 AGA R 2088 GGG G C 2142	GAT D GCA A GAC D	ACT T AGG R AGT S	TCA S 2043 AAG K 2097 CAC H	GCA A A AGA	TCT S TCC S GCA A	TGG W 2052 TCT S 2106 GAG E

FIG.11E

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	-	2169									2196			2205			2114
GAG	CAG	GAC	CAA	GCG	CAA	GGA	AGT	TCC	TCA	CCC	GAG	CCA	GCG	GGA	AGC	CTT	TCC
Ē	Q	D	Q	Α	Q	G	S	S	S	Р	Ε	Р	Α	G	S	Р	S
	2	2223		2	2232		2	2241			2250			2259			2268
GAA	GGG	GAA	GGT	GTC	TCC	ACT	TGG	GAG	TCA	Ш	AAA	AGA	TTA	GTC	ACT	CCA	AGA
E	G	E	G	٧	S	T	W	E	S	F	K	R	L	٧	T	P	R
	2	2277		2	2286		2	2295		,	2304		2	2313		į	2322
AAA	AAA	TCC	AAG	TCA	AAA	CTG	GAA	GAG	AAA	GCC	GAA	GAC	TCT	AGT	GTA	GAG	CAG
K	K	S	K	S	K	L	E	E	K	A	E	D	S	S	٧	E	Q
		2331		2	2340		2	2349		2	2358		2	2367		4	2376
TTG	TCC	ACT	GAG	ATC	GAA	CCG	AGT	AGA	GAA	GAA	TCT	TGG	GTT	TCC	ATT	AAG	AAA
L	S	T	E	I	E	P	S	R	E	E	S	W	٧	S	I	K	K
		2385			2394			2403		2			-	2421		_	2430
ттс	ATC		GGA										-		CAA	_	
TTC  F	ATC			CGG		AAG	AAA 	AGG	GCA	GAC		AAG 	-	GAA	CAA  Q	GCC	
F	ATC I	CCC P 2439	G	CGG R	CGG R R 2448	AAG  K	AAA  K	AGG R R 2457	GCA A	GAC D	GGG  G 2466	AAG  K	CAA Q	GAA E 2475	Q	GCC A	ACT T
F	ATC I	CCC P 2439	G	CGG R	CGG R R 2448	AAG  K	AAA  K	AGG R R 2457	GCA A	GAC D	GGG  G 2466	AAG  K	CAA Q	GAA E 2475	Q	GCC A	ACT T
F	ATC I	P 2439 GAC	G	CGG R R GGG	R R 2448 CCA	AAG  K	AAA K GAG	AGG R 2457 ATA	GCA A AAT	GAC D	GGG  G 2466	AAG  K	CAA Q	GAA E 2475	Q	GCC A	ACT T
F GTG	ATC I GAA E	CCC  P 2439 GAC  D	G TCA	CGG R R GGG	CGG R 2448 CCA P	AAG K GTG V	AAA K GAG E	AGG R 2457 ATA  I	GCA A AAT  N	GAC D GAG  E	GGG  G 2466 GAC  D	AAG K GAC D	CAA Q CCT P	GAA E 2475 AAT N	Q GTC V	GCC A CCA P	ACT T 2484 GCC  A
F GTG	ATC I GAA E	CCC  P 2439 GAC  D	G TCA	CGG R R GGG	CGG R 2448 CCA P	AAG K GTG V	AAA K GAG E	AGG R 2457 ATA  I	GCA A AAT  N	GAC D GAG  E	GGG  G 2466 GAC  D	AAG K GAC D	CAA Q CCT P	GAA E 2475 AAT N	Q GTC V	GCC A CCA P	ACT T 2484 GCC  A
F GTG V	ATC I GAA E	P 2439 GAC D 2493 CCT	G TCA	CGG R GGG G TCT	CGG R 2448 CCA P	AAG K GTG V TAT	AAA K GAG E AAT	AGG R 2457 ATA  I	GCA A  AAT N  GTG	GAC D GAG E GAG	GGG  G 2466 GAC  D	AAG K GAC D GAG	CAA Q CCT P	GAA E 2475 AAT N	Q GTC V	GCC A CCA P	ACT T 2484 GCC  A
F GTG V GTC	ATC I GAA E GTG V	CCC P 2439 GAC D 2493 CCT P	G TCA S CTG	CGG R R GGG TCT S	CGG R R 2448 CCA P 2502 GAG E	AAG K GTG V TAT Y	AAA K GAG E AAT	AGG R 2457 ATA I 2511 GCA A	GCA A  AAT N  GTG	GAC D GAG GAG E GAG GAG	GGG  G 2466 GAC  D 2520 AGG  R	AAG K GAC D GAG	Q 2 CCT P AAG K	GAA E 2475 AAT N 2529 ATG M	Q GTC V GAA	GCC A CCA P GCC	ACT T 2484 GCC  A 2538 CAG  Q
F GTG V GTC	ATC I GAA E GTG	CCC P 2439 GAC D 2493 CCT P	G TCA S CTG	CGG R R GGG TCT S	CGG R R 2448 CCA P 2502 GAG E	AAG K GTG V TAT Y	AAA K GAG E AAT	AGG R 2457 ATA I 2511 GCA A	GCA A  AAT N  GTG	GAC D GAG GAG E GAG GAG	GGG  G 2466 GAC  D 2520 AGG  R	AAG K GAC D GAG	Q 2 CCT P AAG K	GAA E 2475 AAT N 2529 ATG M	Q GTC V GAA	GCC A CCA P GCC	ACT T 2484 GCC  A 2538 CAG  Q

FIG.11F

(25 of 90) 2619 2628 2646 · AGT AAG ACT CTG GTC CAC ACT GTG AGT GTC GCA GTC ATT GAT GGG ACC AGG GCA S K T L V H T V S V A V I D G T R A GTC ACC AGT GTC GAA GAG CGG TCT CCT TCG TGG ATA TCC GCT TCC GTA ACA GAA V T S V E E R S P S W I S A S V T E CCT CTT GAA CAC ACA GCG GGA GAA GCC ATG CCA CCT GTT GAA GAG GTC ACT GAA PLEHTAGEAM PPVEEVTE AAA GAC ATC ATT GCA GAA GAA ACT CCT GTG CTC ACC CAG ACG TTA CCA GAG GGT K D I I A E E T P V L T Q T L P E G AAA GAT GCC CAT GAC GAC ATG GTC ACC AGT GAA GTG GAT TTC ACC TCA GAA GCT K D A H D D M V T S E V D F T S E A GTG ACA GCC ACA GAG ACC TCA GAG GCT CTC CGT ACT GAA GAA GTT ACC GAA GCA V T A T E T S E A L R T E E V T E A TCG GGG GCC GAA GAG ACC ACA GAC ATG GTG TCC GCA GTT TCC CAG CTG ACT GAC SGAEETTDMVSAVSQLTD TCC CCA GAC ACC ACA GAG GAA GCC ACC CCA GTT CAG GAG GTA GAG AGT GGT GTG

FIG.11G

S P D T T E E A T P V Q E V E S G V

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СТА	GAT (	3033 ACA									3060 GCC			3069 CAA			3078 GCA
L	D	T	E	E	E	E	R	Q	T	Q	A	I	L	Q	A	٧	A
GAC	AAG	<b>-</b>	AAA		GAG	TCC	CAG	GTG	CCT	GCA		CAG	ACT		CAG		3132 ACG
D	K	٧	K	E	E	S	Q	V	Р	Α	T	Q	T	٧	Q	R	T
GGG	TCA	3141 AAA									3168 GAG						3186 CTG
G	S	K	Α	L	Ε	K	٧	Ε	Ε	٧	Ε	Ε	D	S	Ε	٧	L
GCT	TCG	3195 GAG									3222 GGA						3240 GGA
A	S	E	K	E	K	D	٧	М	P	K	G	Р	٧	Q	Ε	А	G
GCT	GAG	3195 CAT									3276 CAG			3285 CCA			3294 CTT
		CAT	CTT	GCA	CAG	GGC	TCT		ACT	GGA	CAG		ACT			AGC	
Α	GAG E	CAT  H 3303	CTT L	GCA  A	CAG Q Q 3312	GGC  G	TCT  S	GAG E 83321	ACT T	GGA G	CAG Q 3330	GCT  A	ACT T	CCA P 8339	GAG  E	AGC S	CTT L 3348
Α	GAG E GTT	CAT  H 3303	CTT L GAA	GCA A GTC	CAG Q Q 3312	GGC G GCA	TCT  S GAT	GAG E 3321 GTA	ACT T	GGA G CAT	CAG Q 3330	GCT A GCC	ACT T ACG	P B3339 TGC	GAG  E	AGC S	CTT L 3348
A GAA  E	GAG E GTT V	CAT  H 3303 CCT  P	CTT L GAA	GCA A GTC V	CAG Q 3312 ACG  T	GGC G GCA  A	TCT S GAT D	GAG E 3321 GTA V 3375	ACT T GAC	GGA G CAT  H	CAG Q 3330 GTC  V	GCC A	ACT T ACG	CCA P 3339 TGC  C	GAG E CAG  Q	AGC S GTT V	CTT L 3348 ATC  I
GAA E	GAG E STT V	CAT H 3303 CCT P 3357 CAG	CTT L GAA E CAG	GCA A GTC V CTG	CAG Q 3312 ACG  T 3366 ATG	GGC G GCA A GAA	TCT S GAT D CAG	GAG E 3321 GTA  V 3375 GCC	ACT T  GAC D  GTG	GGA GCC GCC	Q 3330 GTC  V 3384 CCT	GCC A GCC A GAG	ACT T ACG T TCA	CCA P 3339 TGC C 3393 TCC	GAG E CAG Q GAA	AGC S GTT V	CTT L 3348 ATC  I
GAA E AAG	GAG E GTT V CTC	CAT H 3303 CCT P 3357 CAG Q 3411	CTT L GAA E CAG Q	GCA A GTC V CTG	CAG Q 3312 ACG T 3366 ATG  M	GGC GCA A GAA	TCT S GAT D CAG	GAG E 3321 GTA V 3375 GCC  A	ACT T GAC D GTG	GGA GCC A	CAG Q 3330 GTC V 3384 CCT P	GCT A GCC A GAG E	ACT T ACG T TCA S	CCA P 3339 TGC C 3393 TCC S 3447	GAG E CAG Q GAA E	AGC S GTT V ACC	CTT L 3348 ATC I 3402 TTG L

FIG.11H

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GGG	ACA	3465 CAG	CAA	GAT	8474 GAA	ACC	ATT	3483 GAC	AGC	CAG	3492 GAC	AGT	AAA	3501 GCC	ACT	GCA	3510 GCT
G		 Q	 Q					D							 T	 A	
u	'	Ų	Q													_	
CTC	400	3519		CAG	3528 GTC	۸۲۸	CVV.	3537 GAA	GAG	30a	3546 GCT	ΔΟΤ	GCT.	3555 CAG	ΔΔΔ		3564 GAG
<b>GI</b> L	AGG	CAG	1CA			,-											
٧	R	Q	S	Q	٧	T	Ε	E	Ε	Α	Α	T	Α	Q	K	E	Ε
	3	3573		(	3582			3591		3	3600		(	3609		3	3618
CCT	TCG	ACA	CTA	CCT	TAA	AAT	GTT	CCA	GCC	CAG	GAA	GAA	CAT	GGG	GAA	GAA	CCA
P	S	T	L	P	N	N	٧	Р	A	Q	E	E	Н	G	Ε	E	Р
	(	3627			3636		;	3645		;	3654		;	3663			3672
GGA	AGA	GAT	GTT	CTT	GAA	CCT	ACA	CAG	CAA	GAG	CTT	ACT	GCT	GCA	GCC	GTG	CCC
G	R	D	V	L	E	Р	T	Q	Q	E	L	T	А	Α	Α	٧	Р
	,	3681		į	3690									3717			3726
GTT	CTG	GCA	AAG	ACT	GAG	GTG	GGT	CAA	GAG	GGT	GAG	GTT	GAC	TGG	TTG	GAT	GGA
٧	 L	Α	K	T	E	V	G	Q	E	G	E	٧	D	W	L	D	G
		3735			3744			3753		,	3762			3771		;	3780
GAA	AAA	GTC	AAA	GAA	GAA	CAG	GAG	GTG	Ш	GTA	CAC	TCT	GGA	CCC	AAC	AGT	CAA
E	K	٧	K	E	E	Q	E	٧	F	٧	H	S	G	Р	N	S	Q
		3789			3798			3807			3816			3825		;	3834
AAG	GCT	GCT	GAT	GTG	ACA	TAT	GAC	AGT	GAA	GTG	ATG	GGA	GTG	GCC	GGG	TGT	CAG
 К	Α	Α	D	٧	Т	Υ	D	S	E	V	М	G	V	A	G	С	Q
		3843	,					3861						3879			3888
GAA	4.40	CAC	. ACT								CTG	GAG	GAG	GGA	GAG	ATG	GAA
	AAG	i GAC	i AG1	AC I	GAVA	GIG	CAC	MUI	CII	AUC	, ctu	u/ lu	unc	uur	u iu	7110	

FIG.111

## (28 of 90)

ACT		3897 GTT												3933 GTG			3942 GAA
T	D	٧	E	 К	- <b></b> -	 K	 R	 Е		 К	 Р	<del>-</del> -	Q	V	S	E	E
GGT		3951 CAG		ACA										3987 GGG			3996 GTC
G	E	Q	E	T	Α	A	Р	E	Н	Ε	G	T	Y	G	K	Р	V
CTG		1005 CTT												1041 GGA			
L	T	L	D	М	Р	S	S	E	R	G	K	Α	L	G	S	L	G
GGA		4059 CCT												1095 GAG			GTT
G	S	Р	S	L	P	D	Q	D	K	A	G	С	I	E	٧	Q	٧
CAA	AGC	4113 CTG	GAC	ACA	4122 ACA	GTC	ACT	4131 CAA	ACA	GCA	4140 GAA	GCT	GTG	4149 GAA	AAG	GTC	1158 ATA
	AGC	CTG	GAC	ACA	ACA	GTC	ACT	CAA	ACA	GCA	GAA	GCT	GTG	4149 GAA  E	AAG	GTC	ATA
Q	AGC S	CTG  L 4167	GAC D	ACA T	ACA T 4176	GTC  V	ACT T	CAA  Q 4185	ACA T	GCA A	GAA  E 4194	GCT  A	GTG V	GAA	AAG  K	GTC V	ATA I 1212
Q	AGC S ACG	CTG  L 4167	GAC D GTG	ACA T	ACA T 4176 TCA	GTC V GAG	ACT T	CAA Q 4185 GGT	ACA T GAA	GCA A AGT	GAA E 4194 CCA	GCT A GAG	GTG V TGT	GAA E E 4203	AAG K K GGT	GTC V GAC	ATA I 1212
Q GAA  E	AGC S ACG	CTG  L 4167 GTT  V	GAC D GTG V	ACA T ATT	ACA T 4176 TCA  S	GTC V GAG	ACT T ACA T	CAA Q 4185 GGT  G	ACA T GAA	GCA A AGT S	GAA E 4194 CCA  P	GCT A GAG E	GTG V TGT C	GAA E 4203 GTA	AAG K GGT G	GTC V GAC	ATA  I 1212 CAC  H
Q GAA	AGC S ACG T TTA	CTG  L 4167 GTT  V 4221 CCA	GAC D GTG V GCT	ACA T ATT I GAG	ACA T 4176 TCA  S 4230 AAG	GTC V GAG  E	ACA T TCT	CAA Q 4185 GGT  G 4239 GCA	ACA T GAA E ACG	AGT S	GAA E 4194 CCA  P 4248 GGC	GCT A GAG E CAC	TGT C	GAA E 4203 GTA  V	AAG K GGT G	GTC V GAC  A CAG	ATA I 1212 CAC  H 4266 CAT
Q GAA E	AGC S ACG T TTA	CTG  L 4167 GTT  V 4221 CCA  P	GAC D GTG V GCT A	ACA T ATT I GAG	ACA T 4176 TCA  S 4230 AAG  K	GTC V GAG E TCC S	ACT T ACA T TCT	CAA Q 4185 GGT  G 4239 GCA  A	ACA T GAA E ACG T	GCA A AGT S GGT	GAA  E 4194 CCA  P 4248 GGC  G	GCT A GAG E CAC H	TGT C TGG	GAA E 4203 GTA  V 4257 ACT	AAG K GGT G CTT	GTC V GAC  A CAG	ATA I  1212 CAC H  1266 CAT H  1320

FIG.11J

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ATA	GTA	1329 ACT	ССТ	GCT	1338 CCT	GAA	AGC	1347 ACC	СТА	CAT	1356 CCT	GAC	CTA	1365 CAA	GGA	GAA	 1374 ATA
 I	V	T	 Р	Α	 Р	 E	<b>S</b>	T	L	Н	P	D	 L	Q	G	E	I
AGC	4383 C GCA TCC CAG					4401 CGA TCA <b>G</b> AG											
S	A	S	Q	 R	E	R	S	<b>-</b> Е	E	E	D	K	Р	D	A	G	P
GAT	4437 GAT GCT GAC GGC			4446 AAG GAG AG													1482 CCT
D	Α	D	G	K	E	S	T	A	Ī	E	K	٧	L	K	A	E	P
GAG	ATC	1491 CTG	GAA	CTT	1500 GAG	AGT	AAG	4509 AGC	AAC	AAG	1518 ATT	GTG	CTG	1527 AAC	GTC	ATT	1536 CAG
E	 I	 L	 Е	L	<u></u> Е	S	K	\$	N	K	I	٧	L	N	٧	I	Q
ACA	GCC	1545 GTT											GAA				
		GTT	GAC	CAG	TTC	GCA	CGT	ACA	GAA	ACA	GCC	CCC	GAA	ACT	CAT		TAT
т Т	GCC  A	GTT  V 4599	GAC D	CAG Q	TTC  F 4608	GCA  A	CGT R	ACA T 4617	GAA  E	ACA T	GCC  A 4626	CCC P	GAA E	ACT T 4635	CAT H	GCT A	TAT  Y 1644
T	GCC A	GTT V 4599 CAG	GAC D ACC	CAG Q CAG	TTC  F 4608 GTT	GCA -A CCT	CGT R GCA	ACA T 4617 TGC	GAA E E	ACA T	A 4626 GAC	P AGC	GAA E	T 1635 GAG	H CCC	AAC	TAT  Y 1644
T GAT	GCC A TCA	GTT  V 4599 CAG  Q	GAC D ACC T	CAG Q CAG Q	TTC F 4608 GTT V 4662	GCA  A CCT  P	CGT R GCA	ACA T 4617 TGC C 4671	GAA E AGG  R	ACA T CTT	GCC  A 4626 GAC  D	CCC P AGC	GAA E AGG R	ACT T 1635 GAG  E	CCC P	AAC N	TAT  Y 4644 AGA  R
GAT D	GCC A TCA S	GTT V 4599 CAG  Q 4653 ACA	GAC D ACC T AAA	CAG  CAG  Q  ATG	TTC F 4608 GTT  V 4662 AAA	GCA A  CCT P	CGT R GCA A GCC	ACA T 4617 TGC  C 4671 AAG	GAA E  AGG R  ATG	ACA T CTT L AAA	GCC  A 4626 GAC  D 4680 CAC	P AGC S	GAA E AGG  R GTG	ACT T 1635 GAG  E 1689 CCG	CAT H CCC P CAG	AAC AAC N	TAT  Y 1644 AGA  R
T GAT D TGC	GCC A TCA S TGG	GTT V 4599 CAG Q 4653 ACA T T 4707	GAC D ACC T AAA K	CAG  CAG  Q  ATG	TTC F 4608 GTT  V 4662 AAA  K	GCA A CCT P GAT D	GCA A GCC	ACA T 4617 TGC  C 4671 AAG  K	GAA E AGG R ATG	ACA T CTT L AAA	GCC  A 4626 GAC  D 4680 CAC  H	AGC S	GAA E AGG R GTG V	ACT T 1635 GAG  E 1689 CCG  P	CAT H CCC P CAG	AAC AAC N	TAT  Y 1644 AGA  R 4698 AGA  R

FIG.11K

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TTG	CCG	CGC	TTG	CAG	TTG	AAA	GCG	CCG	GTG	TCA	AAG	TAA	GCA	TTG	AGA	4 AGC	TGC
L	P	R	L	Q	L	K	Α	P	V	S	K	*					
CTC	CTC	4815 AAC	CCA	AAG	ATC	CAA	AAG	GAG	CAT	GCT	GCT	GAT	GGC	CCT	CAG	4 CTC 	CAA
AGC	TTA	4869 GCC	CAG	GCA	GAG	GCC	AGT	GCC	TCT	GGA	AAC	CTA	ACC	AAA	GAA	TCC	CCA
GAC	ACC	4923 ACC	GGA	CCA	1932 AAG	CTA	ACC	GAG	GAG	GGC	GAT	CCC	CCA	AAA	GTT	CAG	968 GTC
CAG	GAA	4977 GAA	GAA	ATG	4986 TCT	ACC	AAG	4995 TCA	GTC	AAA	GAG	AAC	AAG	GCC	CAG	GCA	6022 GAA
GAG	GAC	5031 CTG	CAG	GAG	5040 CCA	AAG	GGA	5049 GAC	CTG	GCA	GAA	TCC	TCC	GAT	GII	AGT	IGC
TCA	. TTG	5085 TAC	ATC	TGT	5094 AAG	ACC	AGA	5103 ATG	TGA	AAA	5112 CAA	GTC	ACA	5121 GAA	CAA	GAT	5130 GCT
GCT	GT	5139 F GGG	ACC	TTG	5148 G AGA	CCA	AGA	5157 \ TTT	CAG	AGC	5166 CCA	TGA	CAT	5175 CCA	GAG	AGC	5184 AGG
		5193 C CA	A TGA			}'											

FIG.11L

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SSeCKS 13.2.2



116 -

97.4 -

FIG.12

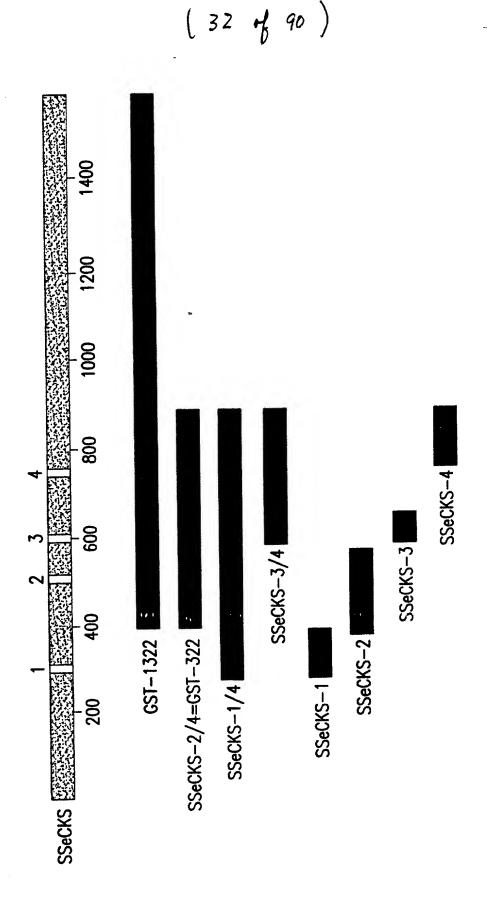


FIG.13A

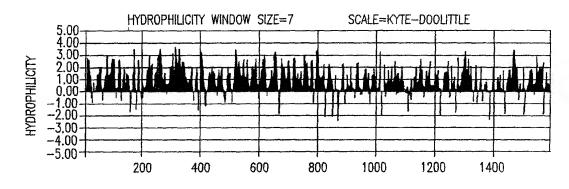


FIG.13B

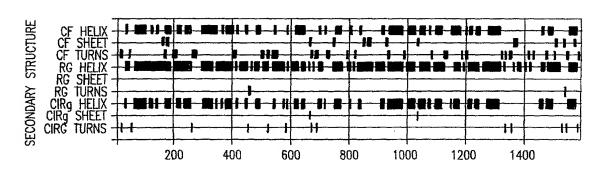
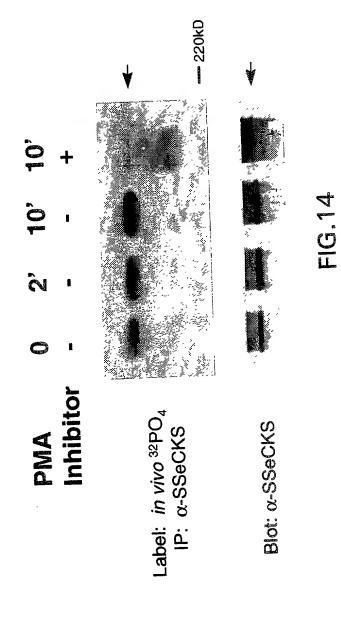
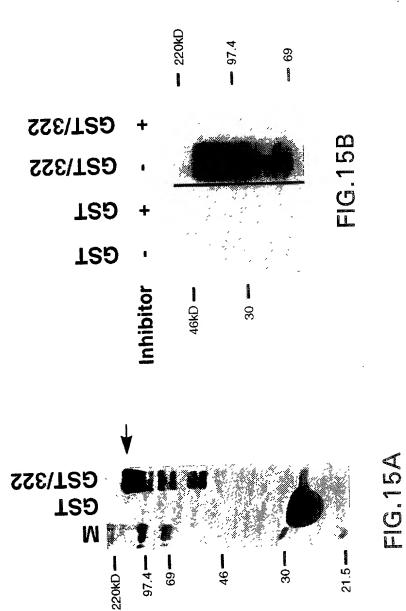


FIG.13C





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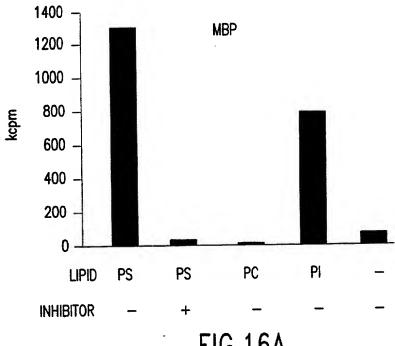
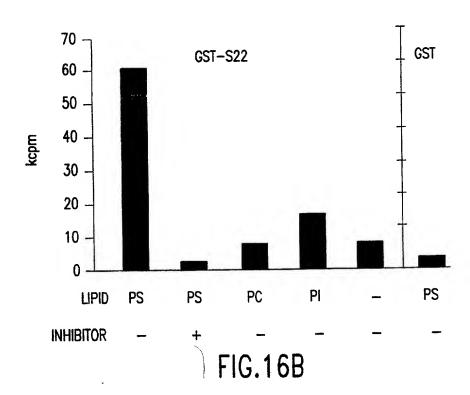
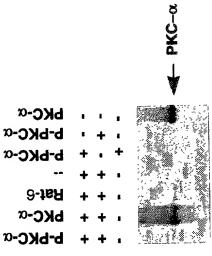


FIG.16A







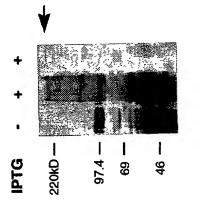
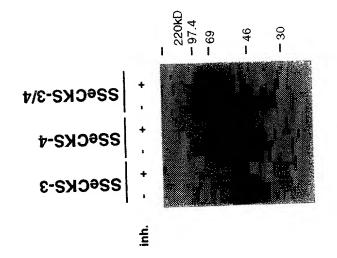


FIG.17A



226CKS-1

226CKS-2

226CK2-1/4

SSeCKS-2/4

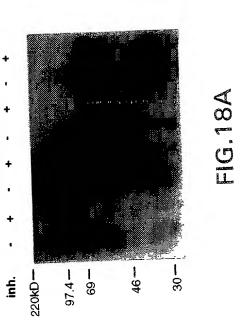
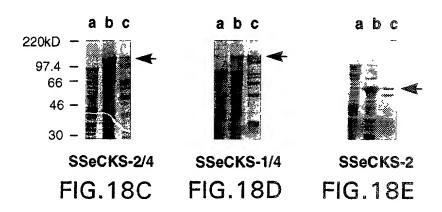


FIG. 18B

#### (39 0/90)





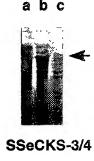


FIG. 181

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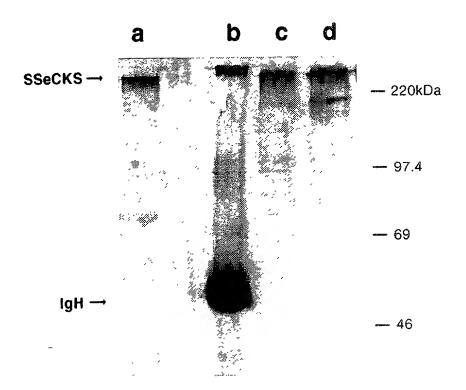
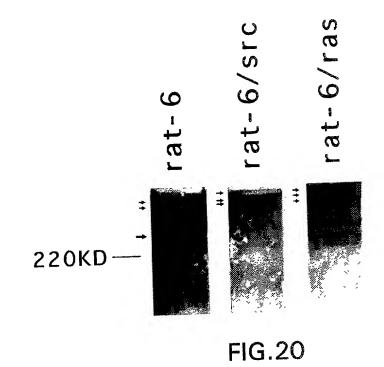


FIG.19



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FIG.21D

FIG.21E

FIG.21F







FIG.21G

FIG.21H

FIG.211

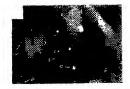


FIG.21J

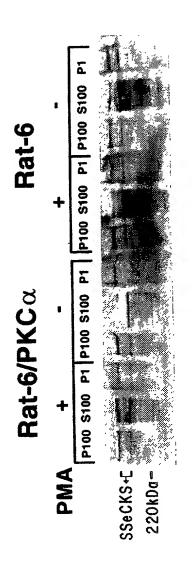


FIG.22

thymus prostate testes testes testes covary colon colon colon

spleen

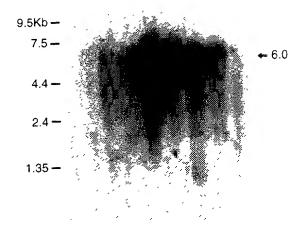
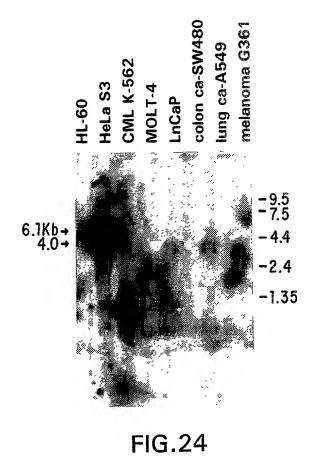


FIG.23A



FIG.23B



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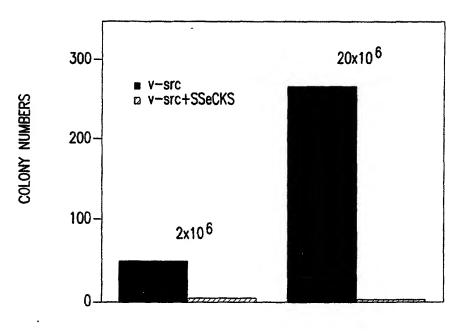


FIG.25A

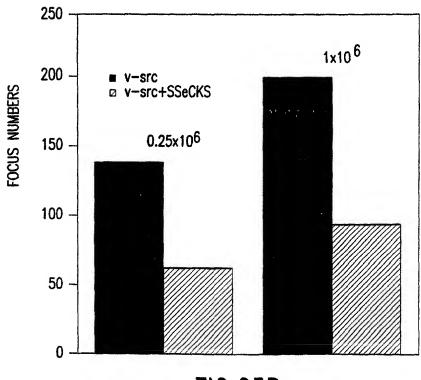


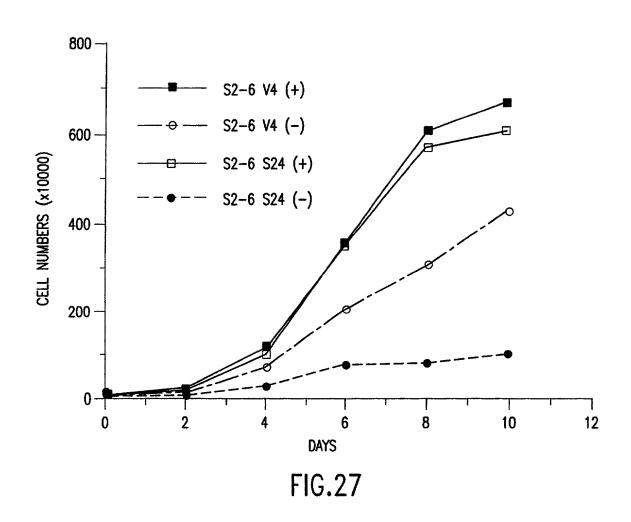
FIG.25B

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		Myr.	Pal.
src	MGSSK <b>S</b> KPKD	+	
yes	<b>MG</b> CIK <b>S</b> KEDK	+	+
SSeCKS	MGAGSSTEQR	+	?
$G_{\alpha t 1}$	<b>MG</b> AGA <b>S</b> AEEK	+	
G <sub>α</sub> i1	MGCTLSAEDK	+	+
GAP-43	M LCCMRRTKQ	-	+
MYRIST. CONCENSUS:	MGXXX <sup>S</sup> / <sub>T</sub>		

FIG.26

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Tet + -



FIG.28

# Swiss 4 mo. weaver 2 wk.

Swiss 2 wk.

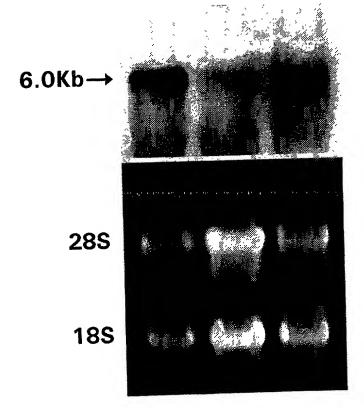


FIG.29

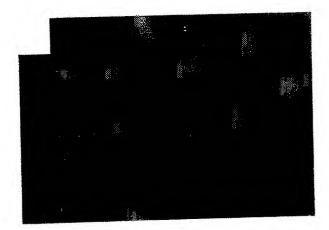


FIG.30A



FIG.30B



FIG.30C

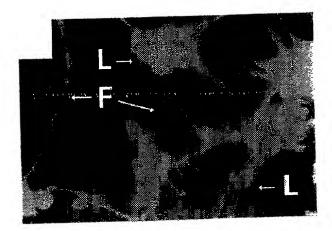


FIG.30D

[53 f 90)

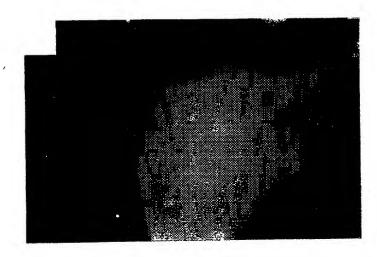


FIG.31A

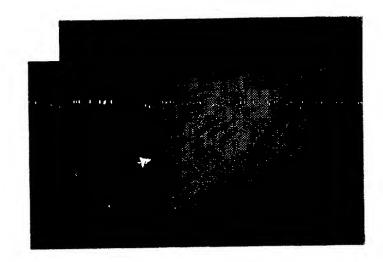


FIG.31B



FIG.31C

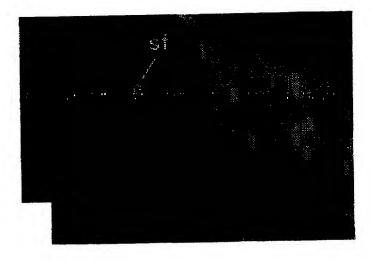
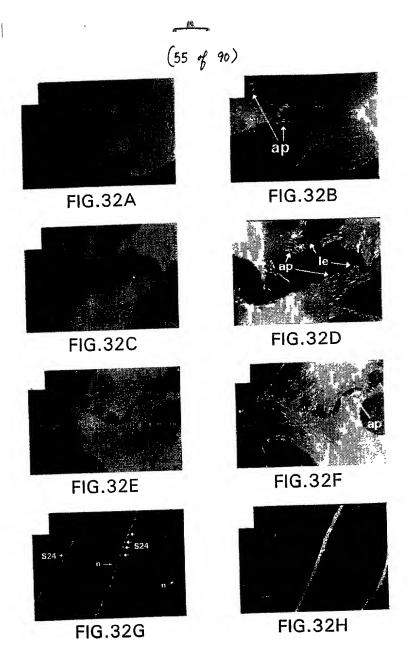


FIG.31D



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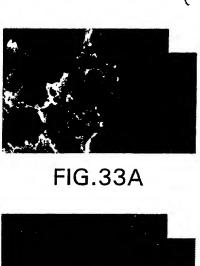










FIG.33D



FIG.33E

FIG.33F

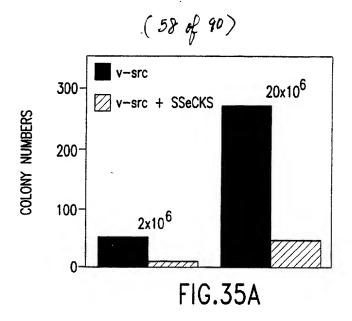


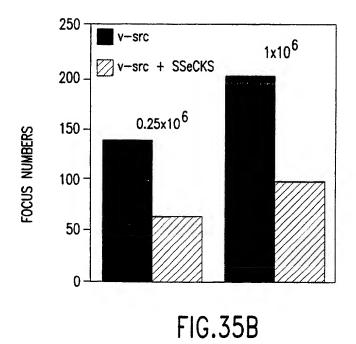


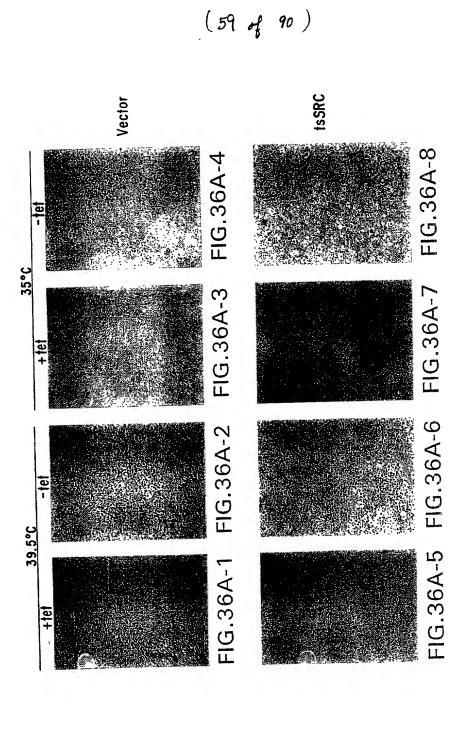
FIG.33G

FIG.33H

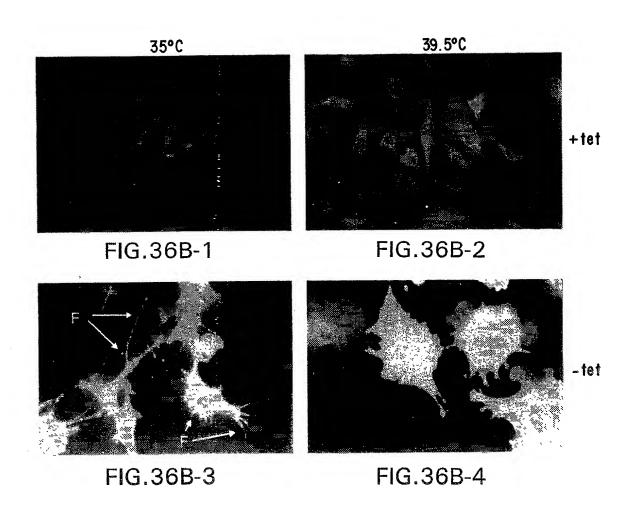
FIG.34







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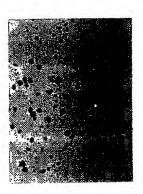


FIG.37A-1



FIG.37A-2

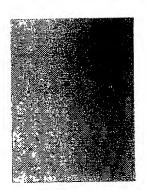


FIG.37A-3

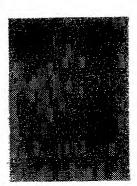


FIG.37A-4

(61 of 90)

		SOFT AGAI	R COLONY FO	ORMATION		
	ts src1	ts src2	ts src3	ts src4	pLJ2	pLJ3
+ tet - tet	2160 60	1640 60	2800 110	1080 35	0 0	0

FIG.37B

#### (63 of 90)

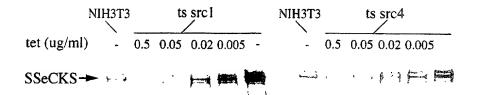


FIG.38A

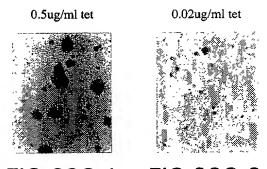


FIG.38C-1 FIG.38C-2

		SOFT AGAR	SOFT AGAR COLONY FORMATION	ORMATION			
		35	35°C			ე <sub>6</sub> €	
tet(ug/ml) 0.5	0.5	0.05	0.02	0.005	0	0.5	0
ts src1	2852	2464	174	51	22	0	0
ts src4	1463	743	29	11	0	0	0

FIG. 38B

### (65 of 90)

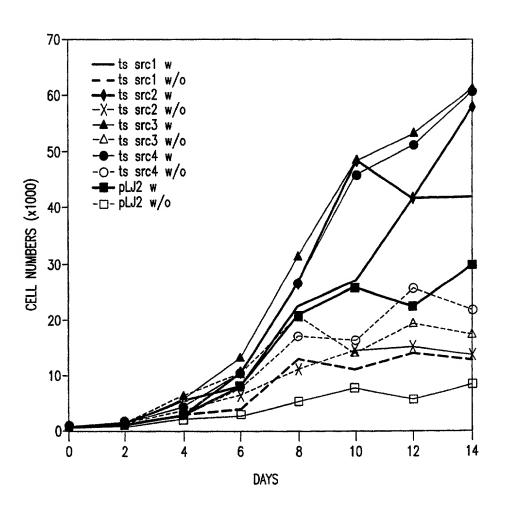


FIG.39A

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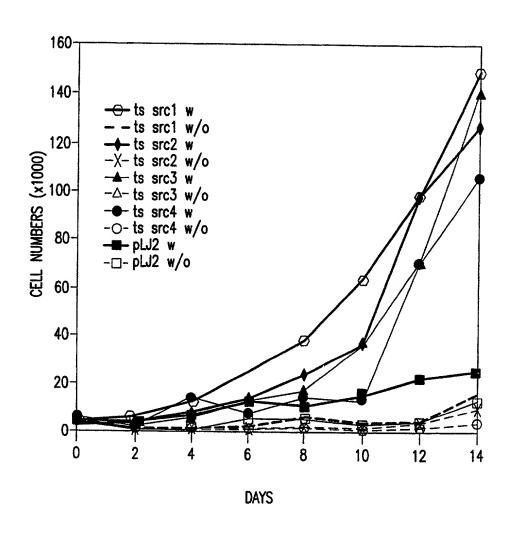
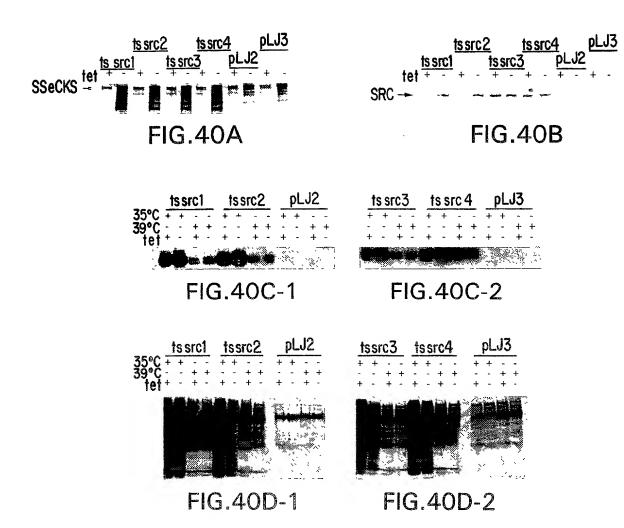
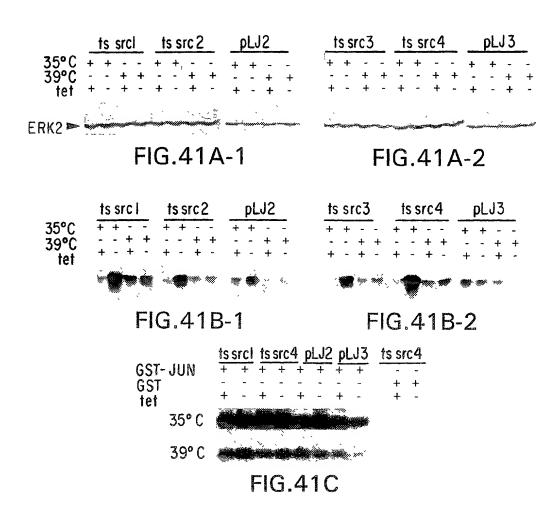


FIG.39B

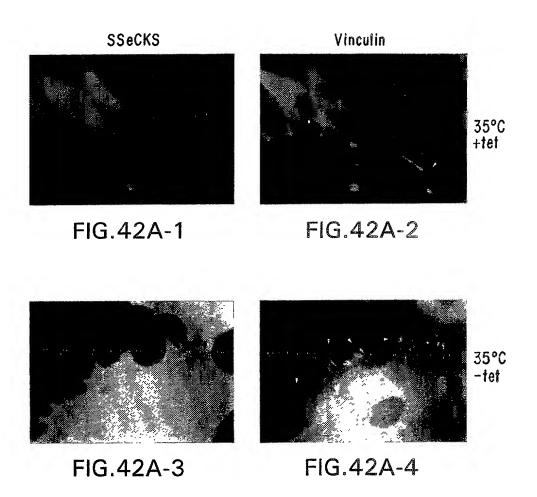
## (67 of 90)



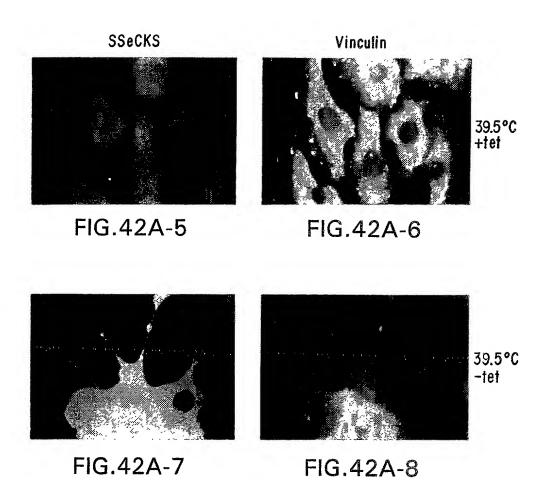
#### (68 of 90)



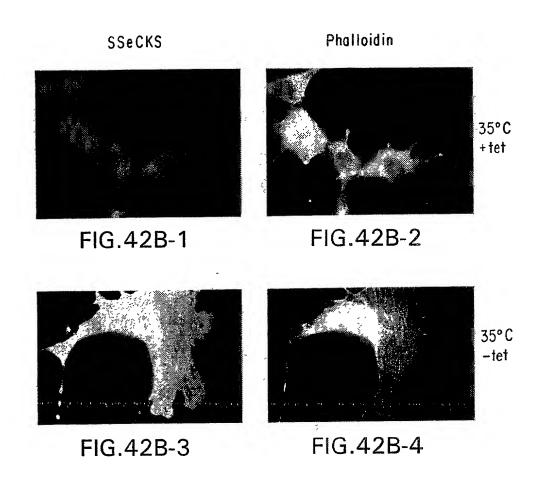
# (69 of 90)



#### (70 of 90)



#### (710f90)



## (72 of 90)

- 5 ...

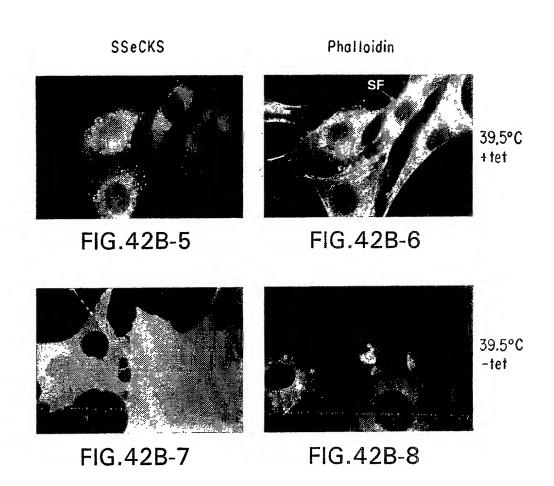
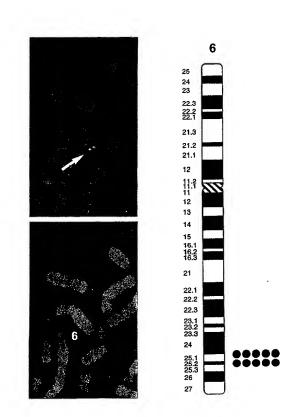


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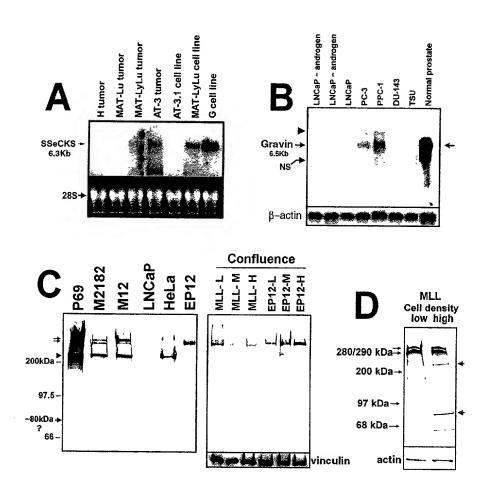


Figure 45
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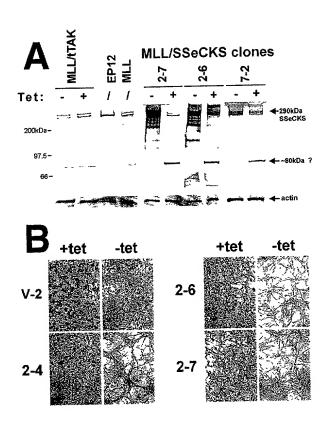
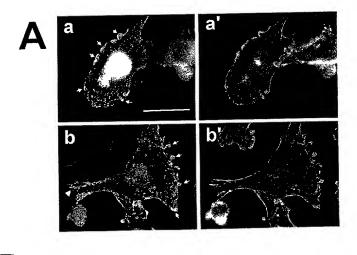
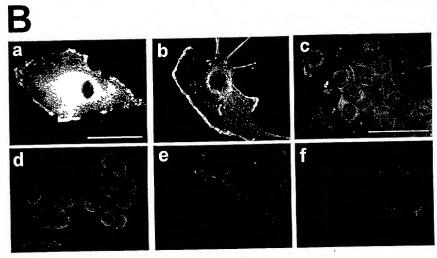


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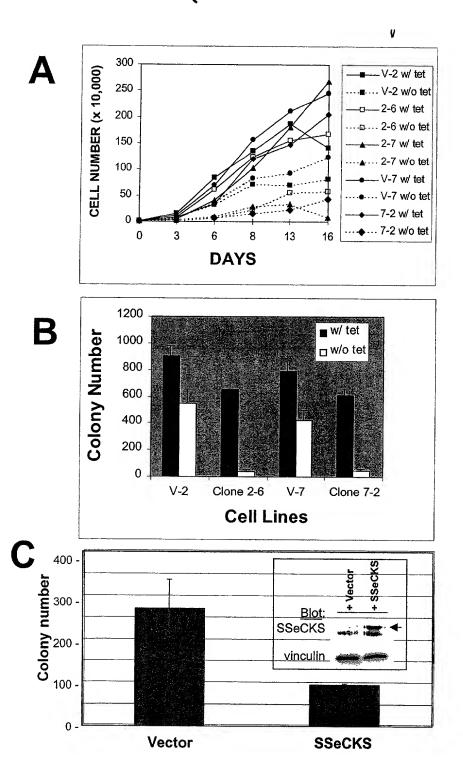


Fig. 44

## (78 of 90)

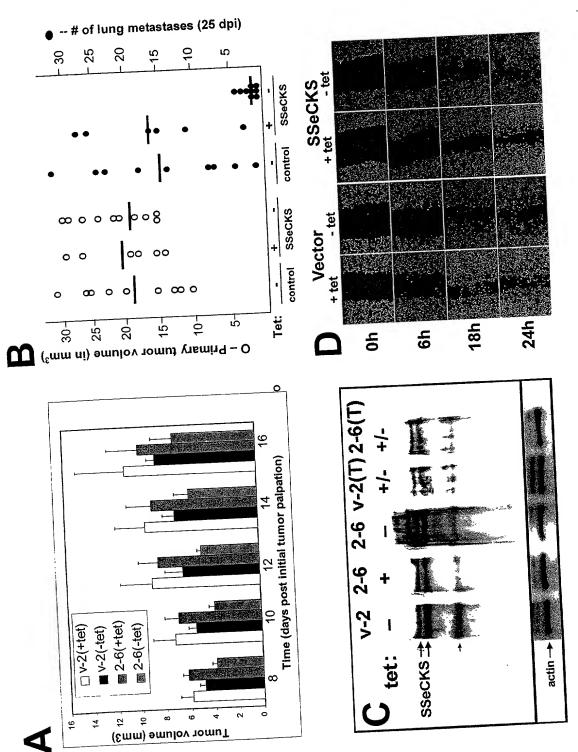


Fig. 48

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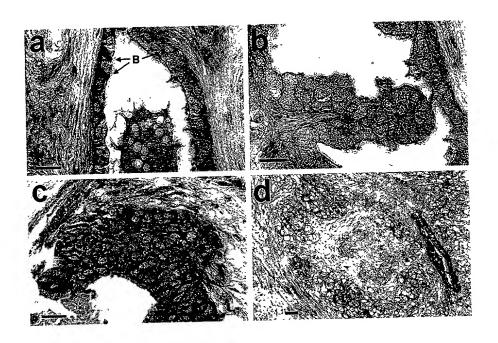


Fig. 49

## (80 of 90)

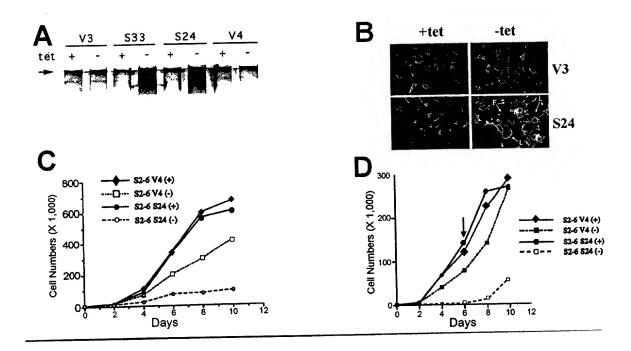


Figure 50

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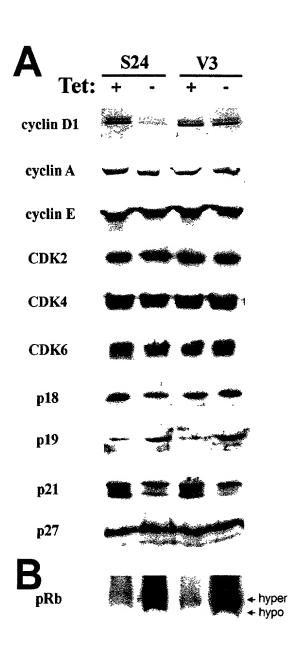


Figure 51

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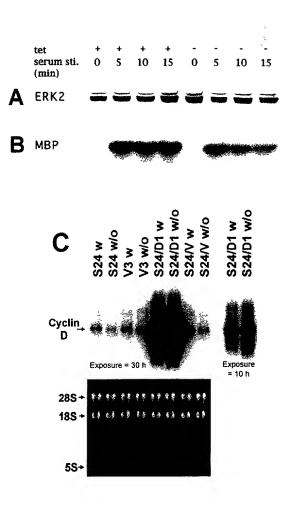


Figure 52

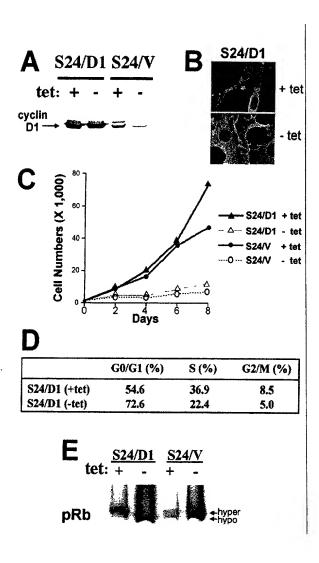


Figure 53

Figure 54

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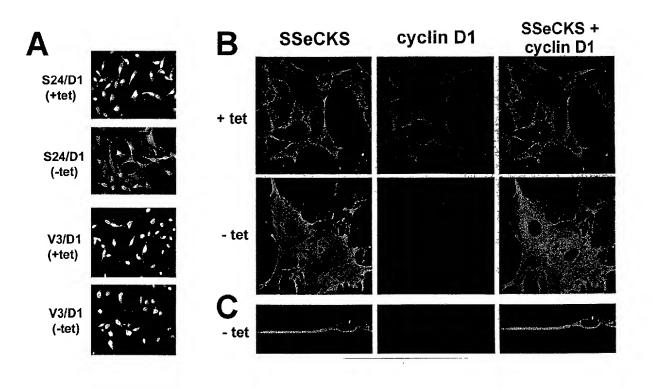


Figure 55

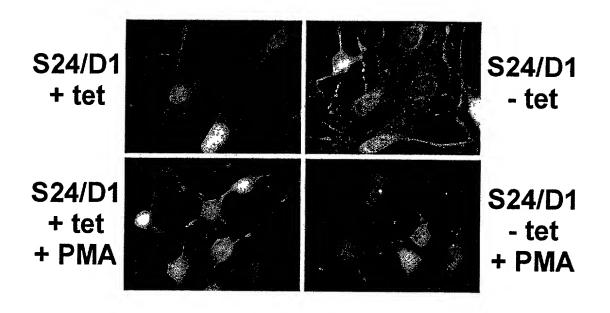


Figure 56

Figure 57
(87 f 90)

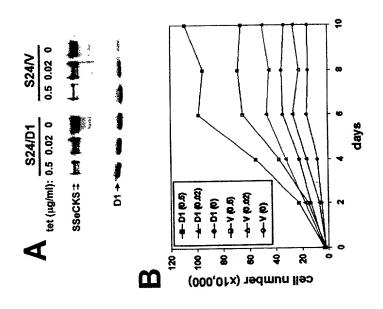
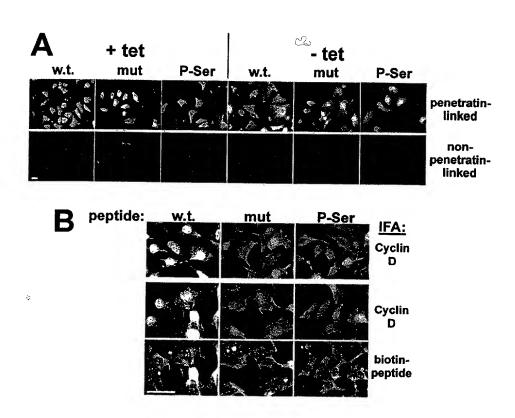


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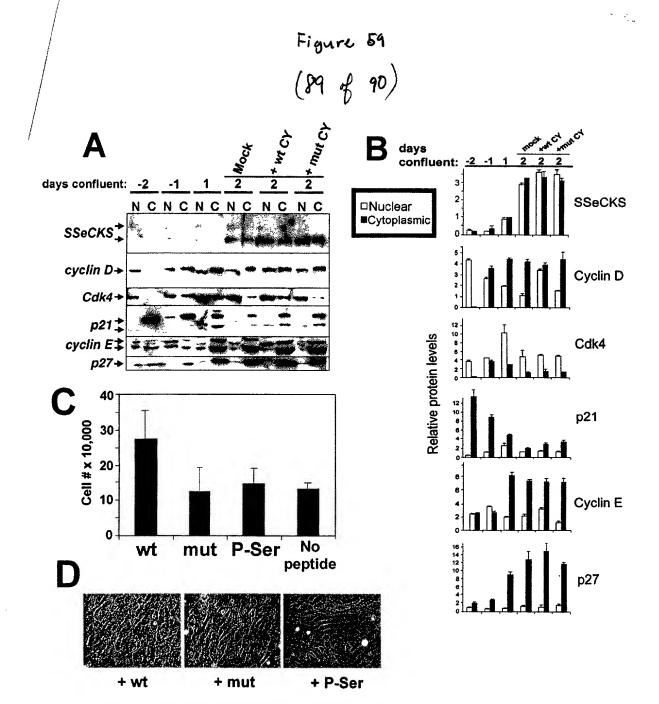
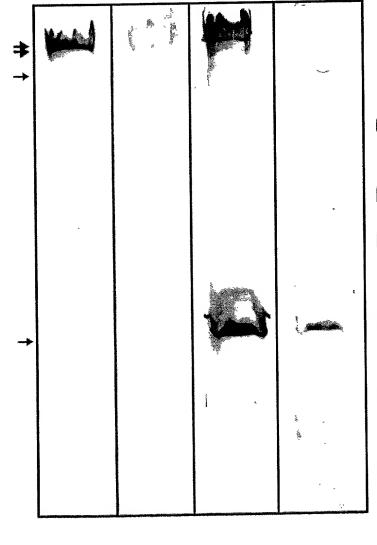


Figure 60 (90 of 90)

94A3 78H11 82B3 31A3



200kDa

97.5

**66** 

**4**4

30

21